STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: $\frac{10/552, 158}{\text{Source:}}$ Date Processed by STIC: $\frac{10/24/2005}{\text{Source:}}$

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual - ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

RAW SEQUENCE LISTING DATE: 10/24/2005
PATENT APPLICATION: US/10/552,158 TIME: 10:47:45

Input Set : A:\Xenon 154.txt

Output Set: N:\CRF4\10242005\J552158.raw

```
4 <110> APPLICANT: Xenon Pharmaceuticals Inc.
      6 <120> TITLE OF INVENTION: Juvenile Hemochromatosis Gene (HFE2A), Expression Products
              and Uses Thereof
      9 <130> FILE REFERENCE: 760050-134
                                                           Does Not Comply
Corrected Diskette Needed

((2) 177, (10)
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/552,158
C--> 12 <141> CURRENT FILING DATE: 2005-10-12
     14 <150> PRIOR APPLICATION NUMBER: 60/462,867
     15 <151> PRIOR FILING DATE: 2003-04-15
     17 <150> PRIOR APPLICATION NUMBER: 60/488,607
     18 <151> PRIOR FILING DATE: 2003-07-18
     20 <150> PRIOR APPLICATION NUMBER: 60/498,458
     21 <151> PRIOR FILING DATE: 2003-08-28
    23 <160> NUMBER OF SEQ ID NOS: (62)
     25 <170> SOFTWARE: PatentIn version 3.0
ERRORED SEQUENCES
E--> 934 <210> SEQ ID NO: 25 (211> 420 (212> PRT (213> Mus musculus (400> 25
E--> 936 <211> LENGTH:
E--> 936 <212> TYPE:
E--> 936 <213> ORGANISM:
E--> 936 <400> SEQUENCE:
     936 Met Gly Gln Ser Pro Ser Pro Arg Ser Pro His Gly Ser Pro Pro Thr
     939 Leu Ser Thr Leu Thr Leu Leu Leu Leu Cys Gly Gln Ala His Ser
     942 Gln Cys Lys Ile Leu Arg Cys Asn Ala Glu Tyr Val Ser Ser Thr Leu
                                     40
     945 Ser Leu Arg Gly Gly Ser Pro Asp Thr Pro Arg Gly Gly Arg
     948 Gly Gly Leu Ala Ser Gly Gly Leu Cys Arg Ala Leu Arg Ser Tyr Ala
     951 Leu Cys Thr Arg Arg Thr Ala Arg Thr Cys Arg Gly Asp Leu Ala Phe
                         85
     954 His Ser Ala Val His Gly Ile Glu Asp Leu Met Ile Gln His Asn Cys
                     100
                                         105
     957 Ser Arg Gln Gly Pro Thr Ala Pro Pro Pro Ala Arg Gly Pro Ala Leu
                115
                                     120
     960 Pro Gly Ala Gly Pro Ala Pro Leu Thr Pro Asp Pro Cys Asp Tyr Glu
                                 135
     963 Ala Arg Phe Ser Arg Leu His Gly Arg Ala Pro Gly Phe Leu His Cys
```

155

150

964 145

Input Set : A:\Xenon 154.txt

966 967	Ala	Ser	Phe	Gly	Asp 165	Pro	His	Val	Arg	Ser 170	Phe	His	Asn	Gln	Phe 175	His
969 970	Thr	Cys	Arg	Val 180	Gln	Gly	Ala	Trp	Pro 185	Leu	Leu	Asp	Asn	Asp 190	Phe	Leu
972 973	Phe	Val	Gln 195	Ala	Thr	Ser	Ser	Pro 200	Val	Ser	Ser	Gly	Ala 205	Asn	Ala	Thr
975 976	Thr	Ile 210	Arg	Lys	Ile	Thr	Ile 215	Ile	Phe	Lys	Asn	Met 220	Gln	Glu	Cys	Ile
979	225		_		-	Gln 230				_	235					240
982		_	_		245	Asn	_	_	_	250		_	_		255	
985				260		Asn			265					270		
988	_		275			Ile		280	_				285			
991		290				Ala	295	_				300				
994	305					Cys 310					315					320
997					325	Asn Glu				330					335	
1000	o			340)				345	5				350)	. Val
100	3	_	355	5				360)				365	;		Leu
100	5	370)				375	·				380)			r Pro
	9 389 1 Ala		е Суа	s Lev	ı Val	390 Pro		ı Lev	ı Ser	. Ala	395 Let		· Val	. Lev	ı Trp	400 Leu
1012		s Phe	e Sei	. Lys	405	5				410)				415	5
1019		L0> S	EQ 1	420 D NC		5										
1019 <211> LENGTH: 422 1020 <212> TYPE: PRT																
1021 <213> ORGANISM: Rattus rattus 1023 <400> SEQUENCE: 26																
	4 Met		_			Arg	Ser	Pro	Ser	Leu 10	Arg	g Ser	Pro	His	Gly 15	/ Ser
102°		Pro	Thr	Let 20	ı Ser	Thr	Leu	Thr	Let 25	Let	Lei	ı Lev	Leu	Cys 30	Gly	/ Gln
1030 1033		a His	Ser 35	Glr	ı Cys	Lys	Ile	Leu 40	a Arg	Cys	Asr	n Ala	Glu 45	Туг	· Val	Ser
1034	1	50					55					60				g Gly
	6 Gly 7 65	/ Gly	Arg	g Gly	/ Gly	7 Pro	Ala	Ser	Gly	gly	7 Let	ı Cys	Arg	Ala	Let	Arg 80

Input Set : A:\Xenon 154.txt

```
1039 Ser Tyr Ala Leu Cys Thr Arg Arg Thr Ala Arg Thr Cys Arg Gly Asp
    1040
    1042 Leu Ala Phe His Ser Ala Val His Gly Ile Glu Asp Leu Met Ile Gln
          100
                                        105
    1045 His Asn Cys Ser Arg Gln Gly Pro Thr Ala Ser Pro Pro Ala Arg Gly
    1046 . 115
                                    120
    1048 Pro Ala Leu Pro Gly Ala Gly Pro Ala Pro Leu Thr Pro Asp Pro Cys
    1049 130
                                135
    1051 Asp Tyr Glu Ala Arg Phe Ser Arg Leu His Gly Arg Thr Pro Gly Phe
                            150
                                                155
    1054 Leu His Cys Ala Ser Phe Gly Asp Pro His Val Arg Ser Phe His Asn
                         165
                                            170
    1057 His Phe His Thr Cys Arg Val Gln Gly Ala Trp Pro Leu Leu Asp Asn
                     180
                                         185
    1060 Asp Phe Leu Phe Val Gln Ala Thr Ser Ser Pro Val Ala Ser Gly Ala
                 195
                                     200
    1063 Asn Ala Thr Thr Ile Arg Lys Ile Thr Ile Ile Phe Lys Asn Met Gln
                                215
    1066 Glu Cys Ile Asp Gln Lys Val Tyr Gln Ala Glu Val Asp Asn Leu Pro
                                                235
                            230
    1069 Ala Ala Phe Glu Asp Gly Ser Val Asn Gly Gly Asp Arg Pro Gly Gly
                        245
                                            250
    1072 Ser Ser Leu Ser Ile Gln Thr Ala Asn Leu Gly Ser His Val Glu Ile
    1073 260
                                        265
    1075 Arg Ala Ala Tyr Ile Gly Thr Thr Ile Ile Val Arg Gln Thr Ala Gly
    1076 275
                                    280
    1078 Gln Leu Ser Phe Ser Ile Arg Val Ala Glu Asp Val Ala Arg Ala Phe
                                 295
    1079 290
    1081 Ser Ala Glu Gln Asp Leu Gln Leu Cys Val Gly Gly Cys Pro Pro Ser
                             310
    1084 Gln Arg Leu Ser Arg Ser Glu Arg Asn Arg Arg Gly Ala Ile Ala Ile
                                            330
                         325
    1087 Asp Thr Ala Arg Arg Leu Cys Lys Glu Gly Leu Pro Val Glu Asp Ala
    1090 Tyr Phe Gln Ser Cys Val Phe Asp Val Ser Val Ser Gly Asp Pro Asn
    1091 355
                                    360
    1093 Phe Thr Val Ala Ala Gln Ser Ala Leu Asp Asp Ala Arg Val Phe Leu
    1094 370
                                375
    1096 Thr Asp Leu Glu Asn Leu His Leu Phe Pro Val Asp Ala Gly Pro Pro
                            390
                                                395
    1097 385
    1099 Leu Ser Pro Ala Thr Cys Leu Val Arg Leu Leu Ser Val Leu Phe Val
                        405
    1102 Leu Trp Phe Cys Ile Gln
    LENGTH: 366
LIU8 <212> TYPE: PRT
1109 <213> ORGANISM: Fugu (2)
1111 <400> SEQUENCE: 27
E--> 1111 <400> SEQUENCE: 27
```

Input Set : A:\Xenon 154.txt

1112		Ser	Cys	Arg	Ile	Leu	Arg	Cys	Asn		Asp	Phe	Val	Ala		Thr
1113 1115		7.00	T 011	Glw	502	802	71-	C111	ת 1 ת	10	C111	C111	ת ד ת	Dro	15	Sor
1116	Бец	voh	neu	20	SCI	SCI	Ата	GIY	25	Gry	GIY	Gry	AIG	30	пец	261
1118	Δra	Glu	Δla		Δen	Δla	Glu	ጥህጕ		Δra	Δla	Len	Hic		Tur	Ser
1119	my	GIU	35	ALG	ASII	nia	Gru	40	Cys	Arg	niu	ыси	45	DCI	- 7 -	DCI
1121	Thr	Cvs		Lvs	Ara	Met	Δla		Pro	Cvs	Ara	Glv		T.e.i	Δla	Tvr
1122		50		-7-			55	5		O _I D		60				-1-
1124	His		Ala	Val	Gln	Glv		Glu	Asp	Leu	Leu		Gln	Tvr	Ara	Cvs
1125						70					75			- 4 -	5	80
1127	Pro	Leu	Ala	Gly	Pro	Thr	Ala	Gln	Pro	Arg	Pro	Leu	Pro	Pro	Leu	Leu
1128				-	85					90					95	
1130	Ser	Gly	Asp	Val	Cys	Leu	Tyr	Asp	Arg	Arg	Leu	Ala	Ala	Ala	Glu	Ala
1131				100					105					110		
1133	Pro	Gln	Pro	Asp	Tyr	Leu	His	Cys	Gly	Val	Phe	Gly	Asp	${\tt Pro}$	His	Ile
1134			115					120					125			
1136	Arg	Thr	Phe	Asn	Asn	Asp	Phe	His	Thr	Cys	Ala	Val	Gln	Gly	Ala	\mathtt{Trp}
1137		130					135					140				
1139	Pro	Leu	Ile	Asp	Asn	_	Phe	Leu	Tyr	Val		Ala	Thr	Ser	Ser	
1140						150				_	155					160
1142	Thr	Arg	Arg	Gly		Gln	Ala	Thr	Met		Thr	Lys	Ile	Thr		Ile
1143		-		m	165	•••	~	7		170	~1			~1	175	~1
1145	vai	Lys	ser	_	Arg	His	Cys	vai	_	GIn	GIn	Leu	Tyr		Ala	GIu
1146	*	7	7	180	D	Mak.	77.	nh a	185	N	a 1	0	77 <u>~</u> 7	190	0	a 1
1148	ьeu	Asp	195	vai	Pro	мет	Ата		Ата	Asp	GIY	ser		vai	ser	GIY
1149 1151	Glu	λνα		G1 v	Gln.	uic	Thr	200	λla	Tla	Thr	Gln.	205 Ser	Dro	Gl v	Ara
1151	Giu	210	Arg	GIY	GIII	птр	215	пеп	нта	116	TIIT	220	Ser	PIO	Gry	AIG
1154	His		Glu	Tle	Ara	Δla		His	Tle	Δla	Thr	_	Δla	Ser	Glv	Gln
1155		nια	Olu	110	nr 9	230	nia	1113	110	niu	235	vui	nια	DCI	Ory	240
1157		Glv	Ara	Ser	Leu		Leu	Ser	۷al	Tvr		Pro	Ara	Ser	Val	
1158		1			245					250			5		255	
1160	Glu	Ala	Phe	Gly	Pro	Glu	Gln	Asp	Leu	Gln	Leu	Cys	Met	Trp	Gly	Cys
1161				260				-	265			_		270	_	_
1163	Pro	Ala	Ser	Gln	Lys	Leu	Ser	Thr	Pro	Pro	Pro	Thr	Ser	Ser	Thr	Phe
1164			275					280					285			
1166	Ser	Ala	Ala	Val	Leu	Ala	His	Cys	Asp	Ala	Leu	Leu	Pro	Val	Arg	Asp
1167		290					295					300				
1169	Val	Tyr	His	Gln	Ala	Cys	Ile	Phe	Asp	Leu	Ile	Thr	Ser	Gly	Asp	Leu
1170				_	_	310	_		_		315		_			320
1172	Asn	Ser	Ser	Gly		Ala	Ile	Ser	Ala		Gln	Asp	Ala	Gln	_	Leu
1173		_	_	_	325				_	330	_	_		_	335	
1175	Ile	Ser	Asp		Lys	Arg	Val	His		Leu	Ser	Pro	Thr		Ala	Ala
1176	a 1 -		~ 3	340		.	~	.	345	.	.	.	-	350		
1178	GIN	arg		Asp	H1S	ьeu	cys		ьeu	ьeu	ьeu	ьeu		ser		
1179	-216	1 - 21	355	. אור	. 20			360					365			
	2 <210> SEQ ID NO: 28 3 <211> LENGTH: 432															
	4 <212> TYPE: PRT															
7104	~~14	11		1 1/1												

Input Set : A:\Xenon 154.txt

Output Set: N:\CRF4\10242005\J552158.raw

1185 <213> ORGANISM: Chicken 1187 <400> SEQUENCE: 28 1188 Met Gly Arg Gly Ala Gly Ser Thr Ala Leu Gly Leu Phe Gln Ile Leu 1191 Pro Val Phe Leu Cys Ile Phe Pro Pro Val Thr Ser Pro Cys Lys Ile 25 1194 Leu Lys Cys Asn Ser Glu Phe Trp Ala Ala Thr Ser Gly Ser His His 40 1197 Leu Gly Ala Glu Glu Thr Pro Glu Phe Cys Thr Ala Leu Arg Ala Tyr 1198 50 55 1200 Ala His Cys Thr Arg Arg Thr Ala Arg Thr Cys Arg Gly Asp Leu Ala 70 1203 Tyr His Ser Ala Val His Gly Ile Asp Asp Leu Met Val Gln His Asn 90 1206 Cys Ser Lys Asp Gly Pro Thr Ser Gln Pro Arg Leu Arg Thr Leu Pro 100 105 1209 Pro Gly Asp Ser Gln Glu Arg Ser Asp Ser Pro Glu Ile Cys His Tyr 1210 115 120 1212 Glu Lys Ser Phe His Lys His Ser Ala Ala Pro Asn Tyr Thr His Cys 130 135 1215 Gly Leu Phe Gly Asp Pro His Leu Arg Thr Phe Thr Asp Thr Phe Gln 150 155 1218 Thr Cys Lys Val Gln Gly Ala Trp Pro Leu Ile Asp Asn Asn Tyr Leu 170 165 1221 Asn Val Gln Val Thr Asn Thr Pro Val Leu Pro Gly Ser Ser Ala Thr 180 185 1224 Ala Thr Ser Lys Leu Thr Ile Ile Phe Lys Ser Phe Gln Glu Cys Val 200 1225 195 1227 Glu Gln Lys Val Tyr Gln Ala Glu Met Asp Glu Leu Pro Ala Ala Phe 1230 Ala Asp Gly Ser Lys Asn Gly Gly Asp Lys His Gly Ala Asn Ser Leu 1231 225 230 235 1233 Lys Ile Thr Glu Lys Val Ser Gly Gln His Ile Glu Ile Gln Ala Lys 250 1236 Tyr Ile Gly Thr Thr Ile Val Val Arg Gln Val Gly Arg Tyr Leu Thr 1237 260 265 1239 Phe Ala Val Arg Met Pro Glu Glu Val Val Asn Ala Val Glu Asp Arg 1240 275 280 1242 Asp Ser Gln Gly Leu Tyr Leu Cys Leu Arg Gly Cys Pro Leu Asn Gln 295 290 300 1245 Gln Ile Asp Phe Gln Thr Phe Arg Leu Ala Gln Ala Ala Glu Gly Arg 310 1248 Ala Arg Arg Lys Gly Pro Ser Leu Pro Ala Pro Pro Glu Ala Phe Thr 325 330 1251 Tyr Glu Ser Ala Thr Ala Lys Cys Arg Glu Lys Leu Pro Val Glu Asp 340 345 1252 1254 Leu Tyr Phe Gln Ser Cys Val Phe Asp Leu Leu Thr Thr Gly Asp Val 360 1257 Asn Phe Met Leu Ala Ala Tyr Tyr Ala Phe Glu Asp Val Lys Met Leu

Input Set : A:\Xenon 154.txt

```
1258
         370
                             375
1260 His Ser Asn Lys Asp Lys Leu His Leu Tyr Glu Arg Thr Arg Ala Leu
                                             395
1261 385
                         390
1263 Ala Pro Gly Asn Ala Ala Pro Ser Glu His Pro Trp Ala Leu Pro Ala
                     405
                                         410
1266 Leu Trp Val Ala Leu Leu Ser Leu Ser Gln Cys Trp Leu Gly Leu Leu
1267
                 420
                                     425
1270 <210> SEQ ID NO: 29
1271 <211> LENGTH: 21
1272 <212> TYPE: DNA
1273 <213> ORGANISM: Artificial
1275 <220> FEATURE:
1276 <223> OTHER INFORMATION: Polynucleotide replication primer
1278 <400> SEQUENCE: 29
1279 tccaagtcag cgactctctc g
                                                                            21
1282 <210> SEQ ID NO: 30
1283 <211> LENGTH: 21
1284 <212> TYPE: DNA
1285 <213> ORGANISM: Artificial
1287 <220> FEATURE:
1288 <223> OTHER INFORMATION: Polynucleotide replication primer
                                          Gire Source of Gal?
1290 <400> SEQUENCE: 30
1291 tccaagtcag tgactctctc g
1294 <210> SEQ ID NO: 31
1295 <211> LENGTH: 21
1296 <212> TYPE: DNA
1297 <213> ORGANISM: Artificial
1299 <220> FEATURE:
1300 <223> OTHER INFORMATION: (Fragment containing polymorphism
1302 <400> SEQUENCE: 31
1303 acctgccgcg gggacctcgc c
                                                                            21
1306 <210> SEQ ID NO: 32
1307 <211> LENGTH: 21
1308 <212> TYPE: DNA
1309 <213> ORGANISM: Artificial
1311 <220> FEATURE:
1312 <223> OTHER INFORMATION: (Fragment containing polymorphism
1314 <400> SEQUENCE: 32
                                                                      _ Same Error
1315 acctgccgcg tggacctcgc c
1318 <210> SEQ ID NO: 33
1319 <211> LENGTH: 21
1320 <212> TYPE: DNA
1321 <213> ORGANISM: Artificial
1323 <220> FEATURE:
1324 <223 > OTHER INFORMATION: Fragment containing polymorphism
1326 <400> SEQUENCE: 33
                                                                            21
1327 gcctgggaaa cctggctgga t
1330 <210> SEQ ID NO: 34
1331 <211> LENGTH: 21
```

Input Set : A:\Xenon 154.txt

```
1332 <212> TYPE: DNA
1333 <213> ORGANISM: Artificial
1335 <220> FEATURE:
1336 <223> OTHER INFORMATION: Fragment containing polymorphism
1338 <400> SEQUENCE: 34
1339 gcctgggaaa gctggctgga t
                                                                            21
1342 <210> SEQ ID NO: 35
1343 <211> LENGTH: 21
1344 <212> TYPE: DNA
1345 <213> ORGANISM: Artificial
1347 <220> FEATURE:
1348 <223> OTHER INFORMATION Fragment containing polymorphism
1350 <400> SEQUENCE: 35
                                                                            21
1351 tecettetgt etttagetea t
1354 <210> SEQ ID NO: 36
                                                                         Same Error
1355 <211> LENGTH: 21
1356 <212> TYPE: DNA
1357 <213> ORGANISM: Artificial
1359 <220> FEATURE:
1360 <223> OTHER INFORMATION:
                             Fragment containing polymorphism
1362 <400> SEQUENCE: 36
1363 tcccttctgt gtttagctca t
                                                                            21
1366 <210> SEQ ID NO: 37
1367 <211> LENGTH: 20
1368 <212> TYPE: DNA
1369 <213> ORGANISM: Artificial
1371 <220> FEATURE:
1372 <223> OTHER INFORMATION: Fragment containing polymorphism
1374 <400> SEQUENCE: 37
                                                                            20
1375 gaggaggagg ccggggtgga
1378 <210> SEQ ID NO: 38
1379 <211> LENGTH: 23
1380 <212> TYPE: DNA
1381 <213> ORGANISM: Artificial
1383 <220> FEATURE:
1384 <223> OTHER INFORMATION: (Fragment containing polymorphism
1386 <400> SEQUENCE: 38
                                                                            23
1387 gaggaggagg aggccggggt gga
1390 <210> SEQ ID NO: 39
1391 <211> LENGTH: 21
1392 <212> TYPE: DNA
1393 <213> ORGANISM: Artificial
1395 <220> FEATURE:
1396 <223> OTHER INFORMATION: Fragment containing polymorphism
1398 <400> SEQUENCE: 39
                                                                            21
1399 gcctccctgc cccggaccct t
1402 <210> SEQ ID NO: 40
1403 <211> LENGTH: 21
1404 <212> TYPE: DNA
```

Input Set : A:\Xenon 154.txt

```
1405 <213> ORGANISM: Artificial
1407 <220> FEATURE:
1408 <223> OTHER INFORMATION: (Fragment containing polymorphism
1410 <400> SEQUENCE: 40
1411 gcctccctgc gccggaccct t
                                                                             21
1414 <210> SEQ ID NO: 41
1415 <211> LENGTH: 21
1416 <212> TYPE: DNA
1417 <213> ORGANISM: Artificial
1419 <220> FEATURE:
1420 <223> OTHER INFORMATION: Fragment containing polymorphism
1422 <400> SEQUENCE: 41
                                                                             21
1423 atggtcgtcc cccggggttc t
1426 <210> SEQ ID NO: 42
1427 <211> LENGTH: 21
1428 <212> TYPE: DNA
1429 <213> ORGANISM: Artificial
1431 <220> FEATURE:
1432 <223> OTHER INFORMATION: Fragment containing polymorphism
1434 <400> SEQUENCE: 42
1435 atggtcgtcc accggggttc t
                                                                             21
1438 <210> SEQ ID NO: 43
1439 <211> LENGTH: 21
1440 <212> TYPE: DNA
1441 <213> ORGANISM: Artificial
1443 <220> FEATURE:
1444 <223> OTHER INFORMATION: Fragment containing polymorphism
1446 <400> SEQUENCE: 43
                                                                             21
1447 cqtcccccqq qqttcttqca t
1450 <210> SEQ ID NO: 44
1451 <211> LENGTH: 21
1452 <212> TYPE: DNA
1453 <213> ORGANISM: Artificial
1455 <220> FEATURE:
1456 <223> OTHER INFORMATION: Fragment containing polymorphism
1458 <400> SEQUENCE: 44
                                                                             21
1459 cgtcccccgg cgttcttgca t
1462 <210> SEQ ID NO: 45
1463 <211> LENGTH: 21
1464 <212> TYPE: DNA
1465 <213> ORGANISM: Artificial
1467 <220> FEATURE:
1468 <223> OTHER INFORMATION: Fragment containing polymorphism
1470 <400> SEQUENCE: 45
1471 gtccaaggag cttggcctct a
                                                                             21
1473 <210> SEQ ID NO: 46
1474 <211> LENGTH: 21
1475 <212> TYPE: DNA
1476 <213> ORGANISM: Artificial
```

Input Set : A:\Xenon 154.txt

```
1478 <220> FEATURE:
1479 <223> OTHER INFORMATION: Fragment containing polymorphism
1481 <400> SEQUENCE: 46
                                                                             21
1482 gtccaaggag attggcctct a
1485 <210> SEQ ID NO: 47
1486 <211> LENGTH: 21
1487 <212> TYPE: DNA
1488 <213> ORGANISM: Artificial
1490 <220> FEATURE:
1491 <223> OTHER INFORMATION: Fragment containing polymorphism
1493 <400> SEQUENCE: 47
1494 cccccatggc gttgggggcc a
                                                                             21
1497 <210> SEQ ID NO: 48
1498 <211> LENGTH: 21
1499 <212> TYPE: DNA
1500 <213> ORGANISM: Artificial
1502 <220> FEATURE:
1503 <223> OTHER INFORMATION: Fragment containing polymorphism
1505 <400> SEQUENCE: 48
1506 ccccatggc tttgggggcc a
                                                                             21
1509 <210> SEQ ID NO: 49
1510 <211> LENGTH: 21
1511 <212> TYPE: DNA
1512 <213> ORGANISM: Artificial
1514 <220> FEATURE:
1515 <223> OTHER INFORMATION: Fragment containing polymorphism
1517 <400> SEQUENCE: 49
1518 taagaacatg caggaatgca t
                                                                             21
1521 <210> SEQ ID NO: 50
1522 <211> LENGTH: 21
1523 <212> TYPE: DNA
1524 <213> ORGANISM: Artificial
1526 <220> FEATURE:
1527 <223> OTHER INFORMATION: Fragment containing polymorphism
1529 <400> SEOUENCE: 50
                                                                             21
1530 taagaacatg aaggaatgca t
1532 <210> SEQ ID NO: 51
1533 <211> LENGTH: 21
1534 <212> TYPE: DNA
1535 <213> ORGANISM: Artificial
1537 <220> FEATURE:
1538 <223> OTHER INFORMATION: Fragment containing polymorphism
1540 <400> SEQUENCE: 51
                                                                             21
1541 gccttctcag ctgaacagga c
1544 <210> SEQ ID NO: 52
1545 <211> LENGTH: 21
1546 <212> TYPE: DNA
1547 <213> ORGANISM: Artificial
1549 <220> FEATURE:
```

Input Set : A:\Xenon 154.txt

Output Set: N:\CRF4\10242005\J552158.raw

1550 <223> OTHER INFORMATION: Fragment containing polymorphism 1552 <400> SEQUENCE: 52 21 1553 gccttctcag gtgaacagga c 1556 <210> SEQ ID NO: 53 1557 <211> LENGTH: 21 1558 <212> TYPE: DNA 1559 <213> ORGANISM: Artificial 1561 <220> FEATURE: 1562 <223> OTHER INFORMATION: Fragment containing polymorphism 1564 <400> SEQUENCE: 53 1565 agatgctggg gttcctcttt c 1568 <210> SEQ ID NO: 54 1569 <211> LENGTH: 21 1570 <212> TYPE: DNA 1571 <213> ORGANISM: Artificial 1573 <220> FEATURE: 1574 <223> OTHER INFORMATION: Fragment containing polymorphism 1576 <400> SEQUENCE: 54 1577 agatgctggg attcctcttt c 21 1580 <210> SEQ ID NO: 55 1581 <211> LENGTH: 20 1582 <212> TYPE: DNA 1583 <213> ORGANISM: Artificial 1585 <220> FEATURE: 1586 <223> OTHER INFORMATION: Forward replication primer 1588 <400> SEQUENCE: 55 1589 cacttgagcc caggaatttg 20 1591 <210> SEQ ID NO: 56 1592 <211> LENGTH: 20 1593 <212> TYPE: DNA 1594 <213> ORGANISM: Artificial 1596 <220> FEATURE: 1597 <223> OTHER INFORMATION: Reverse replication primer 1599 <400> SEQUENCE: 56 1600 gactcactgc agccttgacc 20 1603 <210> SEQ ID NO: 57 1604 <211> LENGTH: 22 1605 <212> TYPE: DNA 1606 <213> ORGANISM: Artificial 1608 <220> FEATURE: 1609 <223> OTHER INFORMATION: Forward replication primer 1611 <400> SEQUENCE: 57 22 1612 gtgtgctaca agtttgccga at 1615 <210> SEQ ID NO: 58 1616 <211> LENGTH: 20 1617 <212> TYPE: DNA 1618 <213> ORGANISM: Artificial 1620 <220> FEATURE: 1621 <223> OTHER INFORMATION: Reverse replication primer

Input Set : A:\Xenon 154.txt

1623	<400> SEQUENCE: 58	
1624	gcttgaaact gggagttgga	20
1627	<210> SEQ ID NO: 59	
1628	<211> LENGTH: 22	
1629	<212> TYPE: DNA .	
1630	<213> ORGANISM: Artificial	
1632	<220> FEATURE:	
1633	<223> OTHER INFORMATION: Forward replication primer	
1635	<400> SEQUENCE: 59	
1636	gggaaatggt cccataattc ct	22
1639	<210> SEQ ID NO: 60	
1640	<211> LENGTH: 19	
1641	<212> TYPE: DNA	
1642	<213> ORGANISM: Artificial	
1644	<220> FEATURE:	
1645	<223> OTHER INFORMATION: Reverse replication primer	
1647	<400> SEQUENCE: 60	
1648	cgccctgcca atatgttct	19
1650	<210> SEQ ID NO: 61	
1651	<211> LENGTH: 22	
1652	<212> TYPE: DNA	
1653	<213> ORGANISM: Artificial	
1655	<220> FEATURE:	
1656	<223> OTHER INFORMATION: Forward replication primer	
1658	<400> SEQUENCE: 61	
1659	ggtacttagc ctcgaaatga ga	22
1662	<210> SEQ ID NO: 62	
1663	<211> LENGTH: 20	
1664	<212> TYPE: DNA	
1665	<213> ORGANISM: Artificial	
1667	<220> FEATURE:	
1668	<223> OTHER INFORMATION: Reverse replication primer	
1670	<400> SEQUENCE: 62	
1671	gtgtcacaca actggttggt	20

Input Set : A:\Xenon 154.txt

Output Set: N:\CRF4\10242005\J552158.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:13,14,15,16,17,18,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45,46 Seq#:47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62 VERIFICATION SUMMARY DATE: 10/24/2005 PATENT APPLICATION: US/10/552,158 TIME: 10:47:47

Input Set : A:\Xenon 154.txt

Output Set: N:\CRF4\10242005\J552158.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:934 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO: 25 <211> 420 <212> PRT <213> Mus musculus <400> 25
L:936 M:282 E: Numeric Field Identifier Missing, <211> is required.
L:936 M:282 E: Numeric Field Identifier Missing, <212> is required.
L:936 M:282 E: Numeric Field Identifier Missing, <213> is required.
L:936 M:200 E: Mandatory Header Field missing, <400> is required.

L:1111 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:27 differs:26 L:23 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (62) Counted (61)